

**IFWO** 

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/734,730

DATE: 09/15/2004 TIME: 10:21:47

Input Set : A:\nidn10314.txt

Output Set: N:\CRF4\09152004\J734730.raw

```
3 <110> APPLICANT: Klaveness, Jo
              Rongved, Pal
      5
              Hogset, Anders
                                                          CHARLE WARE
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              Tolleshaug, Helge
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              Cuthbertson, Alan
      8
              Godal, Aslak
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              Hoff, Lars
     10
              Gogstad, Geir
     11
              Bryn, Klaus
     12
              Naevestad, Anne
     13
              Lovhaug, Dagfinn
     14
              Hellebust, Halldis
     15
              Solbakken, Magne
     17 <120> TITLE OF INVENTION: Diagnostic/Therapeutic Agents
     19 <130> FILE REFERENCE: NIDN-10314
     21 <140> CURRENT APPLICATION NUMBER: 10/734,730
     22 <141> CURRENT FILING DATE: 2003-12-15
     24 <150> PRIOR APPLICATION NUMBER: 09/925,715
     25 <151> PRIOR FILING DATE: 2001-08-10
     27 <150> PRIOR APPLICATION NUMBER: 08/959,206
     28 <151> PRIOR FILING DATE: 1997-10-28
     30 <150> PRIOR APPLICATION NUMBER: 60/049,263
     31 <151> PRIOR FILING DATE: 1997-06-07
     33 <150> PRIOR APPLICATION NUMBER: 60/049,264
     34 <151> PRIOR FILING DATE: 1997-06-06
     36 <150> PRIOR APPLICATION NUMBER: 60/049,266
     37 <151> PRIOR FILING DATE: 1997-06-07
     39 <160> NUMBER OF SEQ ID NOS: 20
     41 <170> SOFTWARE: PatentIn version 3.1
     43 <210> SEQ ID NO: 1
     44 <211> LENGTH: 4
     45 <212> TYPE: PRT
     46 <213> ORGANISM: Artificial Sequence
     48 <220> FEATURE:
     49 <223> OTHER INFORMATION: Description of Artificial Sequence: Heparin sulphate binding
pept
     50
              ide
     52 <400> SEQUENCE: 1
     54 Lys Arg Lys Arg
     55 1
     58 <210> SEQ ID NO: 2
     59 <211> LENGTH: 8
     60 <212> TYPE: PRT
     61 <213> ORGANISM: Artificial Sequence
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Input Set : A:\nidn10314.txt
                     Output Set: N:\CRF4\09152004\J734730.raw
     63 <220> FEATURE:
     64 <223> OTHER INFORMATION: Description of Artificial Sequence: Fibronectin peptide
     66 <400> SEQUENCE: 2
     68 Trp Gln Pro Pro Arg Ala Arg Ile
     72 <210> SEQ ID NO: 3
     73 <211> LENGTH: 13
     74 <212> TYPE: PRT
     75 <213> ORGANISM: Artificial Sequence
     77 <220> FEATURE:
     78 <223> OTHER INFORMATION: Description of Artificial Sequence: Lipopeptide consisting of
hep
              arin sulphate binding peptide and fibronectin peptide
     79
     81 <220> FEATURE:
     82 <221> NAME/KEY: MOD RES
     83 <222> LOCATION: (1)..(1)
     84 <223> OTHER INFORMATION: Dipalmitoyl-lysine
     87 <400> SEQUENCE: 3
     89 Lys Lys Arg Lys Arg Trp Gln Pro Pro Arg Ala Arg Ile
     93 <210> SEQ ID NO: 4
     94 <211> LENGTH: 4
     95 <212> TYPE: PRT
     96 <213> ORGANISM: Artificial Sequence
     98 <220> FEATURE:
     99 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic RGDC sequence
     101 <400> SEQUENCE: 4
     103 Arg Gly Asp Cys
     104 1
     107 <210> SEQ ID NO: 5
     108 <211> LENGTH: 24
     109 <212> TYPE: PRT
     110 <213> ORGANISM: Artificial Sequence
     112 <220> FEATURE:
     113 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic fusion peptide
comp
               rising a PS binding component and a fibronectin peptide sequence
     116 <400> SEQUENCE: 5
     118 Phe Asn Phe Arg Leu Lys Ala Gly Gln Lys Ile Arg Phe Gly Gly
     119 1
     122 Gly Trp Gln Pro Pro Arg Ala Ile
                     2.0
     126 <210> SEQ ID NO: 6
     127 <211> LENGTH: 6
     128 <212> TYPE: PRT
     129 <213> ORGANISM: Artificial Sequence
     131 <220> FEATURE:
    132 <223> OTHER INFORMATION: Description of Artificial Sequence: Biotinylated endothelin-
1 pep
     133
               tide
     135 <220> FEATURE:
     136 <221> NAME/KEY: MOD RES
```

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Input Set : A:\nidn10314.txt
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     137 <222> LOCATION: (1)..(1)
     138 <223> OTHER INFORMATION: Biotin-D-Trp
     141 <400> SEQUENCE: 6
     143 Trp Leu Asp Ile Ile Trp
     144 1
     147 <210> SEQ ID NO: 7
     148 <211> LENGTH: 10
     149 <212> TYPE: PRT
     150 <213> ORGANISM: Artificial Sequence
     152 <220> FEATURE:
     153 <223> OTHER INFORMATION: Description of Artificial Sequence: Biotinylated fibrin-
antipolym
     154
               erant peptide
     156 <220> FEATURE:
     157 <221> NAME/KEY: MOD RES
     158 <222> LOCATION: (1)..(1)
     159 <223> OTHER INFORMATION: Biotin-Gly
     162 <220> FEATURE:
     163 <221> NAME/KEY: MOD RES
     164 <222> LOCATION: (10)..(10)
     165 <223> OTHER INFORMATION: AMIDATION
     168 <400> SEQUENCE: 7
     170 Gly Pro Arg Pro Pro Glu Arg His Gln Ser
     171 1
     174 <210> SEQ ID NO: 8
     175 <211> LENGTH: 6
     176 <212> TYPE: PRT
     177 <213> ORGANISM: Artificial Sequence
     179 <220> FEATURE:
     180 <223> OTHER INFORMATION: Description of Artificial Sequence: Lipopeptide
     182 <220> FEATURE:
     183 <221> NAME/KEY: MOD RES
     184 <222> LOCATION: (5)..(5)
     185 <223> OTHER INFORMATION: Biotinylated-lys
     188 <220> FEATURE:
     189 <221> NAME/KEY: MOD RES
     190 <222> LOCATION: (1)..(1)
     191 <223> OTHER INFORMATION: Dipalmitoyl-lysine
     194 <400> SEQUENCE: 8
     196 Lys Trp Lys Lys Gly
     197 1
     200 <210> SEQ ID NO: 9
     201 <211> LENGTH: 25
     202 <212> TYPE: DNA
     203 <213> ORGANISM: Artificial Sequence
     205 <220> FEATURE:
     206 <223> OTHER INFORMATION: Description of Artificial Sequence: Biotinylated synthetic
oligon
     207
               ucleotide
     209 <220> FEATURE:
     210 <221> NAME/KEY: misc feature
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                                                              TIME: 10:21:47
                     Input Set : A:\nidn10314.txt
                     Output Set: N:\CRF4\09152004\J734730.raw
     211 <222> LOCATION: (1)..(1)
     212 <223> OTHER INFORMATION: Biotinylated
     215 <400> SEQUENCE: 9
     216 gaaaggtagt ggggtcgtgt gccgg
                                                                                 25
     219 <210> SEQ ID NO: 10
     220 <211> LENGTH: 25
     221 <212> TYPE: DNA
     222 <213> ORGANISM: Artificial Sequence
     224 <220> FEATURE:
     225 <223> OTHER INFORMATION: Description of Artificial Sequence: Biotinylated synthetic
oligon
     226
               ucleotide
     228 <220> FEATURE:
     229 <221> NAME/KEY: misc feature
     230 <222> LOCATION: (1)..(1)
     231 <223> OTHER INFORMATION: Biotinylated
     234 <400> SEQUENCE: 10
     235 ggcgctgatg atgttgttga ttctt
                                                                                 25
     238 <210> SEQ ID NO: 11
     239 <211> LENGTH: 5
     240 <212> TYPE: PRT
     241 <213> ORGANISM: Artificial Sequence
     243 <220> FEATURE:
     244 <223> OTHER INFORMATION: Description of Artificial Sequence: Lipopeptide containing
the RGD
     245
               sequence and a fluorescein reporter group
     247 <220> FEATURE:
     248 <221> NAME/KEY: MOD RES
     249 <222> LOCATION: (1)..(1)
     250 <223> OTHER INFORMATION: Dipalmitoyl-lys
     253 <220> FEATURE:
     254 <221> NAME/KEY: MOD RES
     255 <222> LOCATION: (4)..(4)
     256 <223> OTHER INFORMATION: Acetyl-RGD-K-fluorescein side chain
     259 <400> SEQUENCE: 11
     261 Lys Lys Lys Gly
     262 1
     265 <210> SEQ ID NO: 12
     266 <211> LENGTH: 18
     267 <212> TYPE: PRT
     268 <213> ORGANISM: Artificial Sequence
     270 <220> FEATURE:
     271 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic endothelial
cell bi
     272
               nding lipopeptide
     274 <220> FEATURE:
     275 <221> NAME/KEY: MOD RES
     276 <222> LOCATION: (1)..(1)
     277 <223> OTHER INFORMATION: 2-n-hexadecylstearyl-lysine
     280 <220> FEATURE:
     281 <221> NAME/KEY: MOD_RES
     282 <222> LOCATION: (18)..(18)
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Input Set : A:\nidn10314.txt
                     Output Set: N:\CRF4\09152004\J734730.raw
     283 <223> OTHER INFORMATION: AMIDATION
     286 <400> SEQUENCE: 12
     288 Lys Leu Ala Leu Lys Leu Ala Leu Lys Ala Leu Lys Ala Ala Leu Lys
     289 1
     292 Leu Ala
     296 <210> SEO ID NO: 13
     297 <211> LENGTH: 5
     298 <212> TYPE: PRT
     299 <213> ORGANISM: Artificial Sequence
     301 <220> FEATURE:
     302 <223> OTHER INFORMATION: Description of Artificial Sequence: Thiol functionalised
lipid mo
     303
               lecule
     305 <220> FEATURE:
     306 <221> NAME/KEY: MOD_RES
     307 <222> LOCATION: (1)..(1)
     308 <223> OTHER INFORMATION: Dipalmitoyl-lysine
     311 <220> FEATURE:
     312 <221> NAME/KEY: MISC_FEATURE
     313 <222> LOCATION: (4)..(4)
     314 <223> OTHER INFORMATION: Acp
     317 <400> SEQUENCE: 13
W--> 319 Lys Lys Lys Xaa Cys
     320 1
     323 <210> SEO ID NO: 14
     324 <211> LENGTH: 4
     325 <212> TYPE: PRT
     326 <213> ORGANISM: Artificial Sequence
     328 <220> FEATURE:
     329 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic lipopeptide
functio
     330
               nalised with captopril
     332 <220> FEATURE:
     333 <221> NAME/KEY: MOD RES
     334 <222> LOCATION: (1)..(1)
     335 <223> OTHER INFORMATION: Dipalmitoyl-lysine
     338 <220> FEATURE:
     339 <221> NAME/KEY: MOD RES
     340 <222> LOCATION: (4)..(4)
     341 <223> OTHER INFORMATION: AMIDATION
     344 <220> FEATURE:
     345 <221> NAME/KEY: MOD RES
     346 <222> LOCATION: (4)..(4)
     347 <223> OTHER INFORMATION: Amide-linked via side chain to captopril
     350 <400> SEQUENCE: 14
     352 Lys Lys Lys
     353 1
     356 <210> SEQ ID NO: 15
     357 <211> LENGTH: 13
     358 <212> TYPE: PRT
     359 <213> ORGANISM: Artificial Sequence
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/734,730

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/734,730

DATE: 09/15/2004 TIME: 10:21:48

Input Set : A:\nidn10314.txt

Output Set: N:\CRF4\09152004\J734730.raw

## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:13; Xaa Pos. 4
Seq#:15; Xaa Pos. 4

VERIFICATION SUMMARY

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,730 TIME: 10:21:48

Input Set : A:\nidn10314.txt

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L:319 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0 L:391 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0